

XM048113	1	-----
ANH401	1	MAAVSLRLGDLVWGKLGRYPPWPGKIVNPPKDLKKPRGKKCFVKKFFGTEDHAWIKVEQL
AF326966	1	MAAVSLRLGDLVWGKLGRYPPWPGKIVNPPKDLKKPRGKKCFVKKFFGTEDHAWIKVEQL
XM048113	1	-----
ANH401	61	KPYHAHKEEMIKINKGKRFQQAVDAVEEFLRRAGKGDQTSSHNSSDDKNRRNSSEERSRP
AF326966	61	KPYHAHKEEMIKINKGKRFQQAVDAVEEFLRRAGKGDQTSSHNSSDDKNRRNSSEERSRP
XM048113	1	-----
ANH401	121	NSGDEKRKLSLSEGVKVKKNMGEGKKRVSSGSSSERGSKSPLKRAQEQQSPRKRGRPPKDEKD
AF326966	121	NSGDEKRKLSLSEGVKVKKNMGEGKKRVSSGSSSERGSKSPLKRAQEQQSPRKRGRPPKDEKD
XM048113	1	-----
ANH401	181	LTIPESSTVKGMMAGPMAAFKWQPTASEPVKDADPHFHHFLLSQTEKPAVCYQAITKKLK
AF326966	181	LTIPESSTVKGMMAGPMAAFKWQPTASEPVKDADPHFHHFLLSQTEKPAVCYQAITKKLK
XM048113	1	-----MSGIVSNLLKMGHTVTVWNRTA
ANH401	241	ICBEETGSTSIQAADSTAVNGSITPTDKKIGFLGLGLMSGIVSNLLKMGHTVTVWNRTA
AF326966	241	ICBEETGSTSIQAADSTAVNGSITPTDKKIGFLGLGLMSGIVSNLLKMGHTVTVWNRTA
XM048113	24	EKCDLFIQEGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGP SGVLQGIRPGKCYVDM
ANH401	301	EKCDLFIQEGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGP SGVLQGIRPGKCYVDM
AF326966	301	EK-----EGARLGRTPAEVVSTCDITFACVSDPKAAKDLVLGP SGVLQGIRPGKCYVDM
XM048113	84	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
ANH401	361	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
AF326966	355	STVDADTVTELAQVIVSRGGRFLEAPVSGNQQLSNDGMLVILAAGDRGLYEDCSSCFQAM
XM048113	144	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAHVTGQSQQTLTDILNQGLASI
ANH401	421	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAHVTGQSQQTLTDILNQGLASI
AF326966	415	GKTSFFLGEVGNAAKMMLIVNMVQGSFMATIAEGLTLAQTGQSQQTLTDILNQGLASI
XM048113	204	FLDQKCQNILOGNFKPDEFYLYIQKDLRLAIALGDAVNHPTPMAAAAANEVYKRAKALDQS
ANH401	481	FLDQKCQNILOGNFKPDEFYLYIQKDLRLAIALGDAVNHPTPMAAAAANEVYKRAKALDQS
AF326966	475	FLDQKCQNILOGNFKPDEFYLYIQKDLRLAIALGDAVNHPTPMAAAAANEVYKRAKALDQS
XM048113	264	DNDMSAVYRAYIH
ANH401	541	DNDMSAVYRAYIH
AF326966	535	DNDMSAVYRAYIH

Fig. 1